## SEQUENCE LISTING

<110> Bristol-Myers Squibb Company

<120> POLYNUCLEOTIDE ENCODING A NOVEL HUMAN POTASSIUM CHANNEL BETA-SUBUNIT, K+betaM2

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Pro Glu Val Val Glu Leu Asn Val Gly Gly Gln Val Tyr Phe Thr Arg

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		_	_	_	acg Thr	_		_		_	-	~		_		727
				_	aga Arg	_			-		-			_	-	775
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				_	ctc Leu		_	_	_	_						1015
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-	_	_		_	aga Arg				-						_	1207
	_	_		-	ttt Phe	_	_	_			_				_	1255
					tcg Ser			_								1303
gat	gac	aag	atc	tgg	tca	agc	tac	act	gaa	tat	gtc	ttc	tac	cgt	gag	1351

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aca tot age tge gae age cag tot gag gee age tot co Thr Ser Ser Cys Asp Ser Gln Ser Glu Ala Ser Ser P: 315	
gtc atc tgt ggt ccc gtg aca cgc cag acc aac atc co Val Ile Cys Gly Pro Val Thr Arg Gln Thr Asn Ile G 330 335 3	
cgt ccc atc aag aag ggc cct gtc cag ctg atc caa ca Arg Pro Ile Lys Lys Gly Pro Val Gln Leu Ile Gln G 345 350 355	
cgg cgg aaa agc gac tta ctc cgg att ctg act tca g Arg Arg Lys Ser Asp Leu Leu Arg Ile Leu Thr Ser G 360 365 370	
tcg aac atg agc agc aaa aaa aaa gct gtt aaa gaa a Ser Asn Met Ser Ser Lys Lys Lys Ala Val Lys Glu L 380 385	
gag gag gag ctg gag aaa tgt atc cag gat ttc cta a Glu Glu Glu Leu Glu Lys Cys Ile Gln Asp Phe Leu L 395 400	
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cattttgaaa ttaacctcat aaaaggaatt catattttaa aggaa	aaaaa tacaactaat 1898
gatgcacatt tettagaaca caatagteea ttgatataet aetge	ctact ttacctagtt 1958
caccttaaca tgtaaatcca cagggtagat ttctttctag atgtg	gaagt acaagaaaat 2018
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Ser Ala Val Pro Asn Ser Phe Pro Glu Val Val Glu Leu Asn Val Gly 25

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Ser Leu Leu Trp Lys Met Phe Ser Pro Lys Arg Asp Thr Ala Asn Asp 50 55 60

Leu Ala Lys Asp Ser Lys Gly Arg Phe Phe Ile Asp Arg Asp Gly Phe 65 70 75 80

Leu Phe Arg Tyr Ile Leu Asp Tyr Leu Arg Asp Arg Gln Val Val Leu 85 90 95

Pro Asp His Phe Pro Glu Lys Gly Arg Leu Lys Arg Glu Ala Glu Tyr 100 105 110

Phe Gln Leu Pro Asp Leu Val Lys Leu Leu Thr Pro Asp Glu Ile Lys 115 120 125

Gln Ser Pro Asp Glu Phe Cys His Ser Asp Phe Glu Asp Ala Ser Gln 130 135 140

Arg Lys Trp Gly Phe Ile Thr Val Gly Tyr Arg Gly Ser Cys Thr Leu 165 170 175

Gly Arg Glu Gly Gln Ala Asp Ala Lys Phe Arg Arg Val Pro Arg Ile 180 185 190

Leu Val Cys Gly Arg Ile Ser Leu Ala Lys Glu Val Phe Gly Glu Thr 195 200 205

Leu Asn Glu Ser Arg Asp Pro Asp Arg Ala Pro Glu Arg Tyr Thr Ser 210 215 220

Arg Phe Tyr Leu Lys Phe Lys His Leu Glu Arg Ala Phe Asp Met Leu 225 230 235 240

Ser Glu Cys Gly Phe His Met Val Ala Cys Asn Ser Ser Val Thr Ala  $245 \hspace{1.5cm} 250 \hspace{1.5cm} 255$ 

Ser	Phe	Ile	Asn 260	Gln	Tyr	Thr	Asp	Asp 265	Lys	Ile	Trp	Ser	Ser 270	Tyr	Thr		
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Asp	Cys 290	Cys	Cys	Lys	Asn	Gly 295	Lys	Gly	Asp	Lys	Glu 300	Gly	Glu	Ser	Gly		
Thr 305	Ser	Cys	Asn	Asp	Leu 310	Ser	Thr	Ser	Ser	Cys 315	Asp	Ser	Gln	Ser	Glu 320		
Ala	Ser	Ser	Pro	Gln 325	Glu	Thr	Val	Ile	Cys 330	Gly	Pro	Val	Thr	Arg 335	Gln		
Thr	Asn	Ile	Gln 340	Thr	Leu	Asp	Arg	Pro 345	Ile	Lys	Lys	Gly	Pro 350	Val	Gln		
Leu	Ile	Gln 355	Gln	Ser	Glu	Met	Arg 360	Arg	Lys	Ser	Asp	Leu 365	Leu	Arg	Ile		
Leu	Thr 370	Ser	Gly	Ser	Arg	Glu 375	Ser	Asn	Met	Ser	Ser 380	Lys	Lys	Lys	Ala		
Val 385	Lys	Glu	Lys	Leu	Ser 390	Ile	Glu	Glu	Glu	Leu 395	Glu	Lys	Cys	Ile	Gln 400		
Asp	Phe	Leu	Lys	Lys 405	Lys	Ile	Pro	Asp	Arg 410	Phe	Pro	Glu	Arg	Lys 415	His		
Pro	Trp	Gln	Ser 420	Glu	Leu	Leu	Arg	Lys 425	Tyr	His	Leu						
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caa	tgąc	tct	gagt	ggaa	ac t	gtag	tcgt	t at	tatc	ctcg	aga	acaa	ggg	tccg	cagttc	3	180
cca	actc	ctt	ccct	gagg	tg g	taga	gctg	a at	gtcg	gggg	tca	agtt	tat	ttta	ctcgcc	2	240

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Tyr Pro Asp Ser Met Leu Gly Ala Met Phe Gly Gly Asp Phe Pro Thr 35 40 45

Ala Arg Asp Pro Gln Gly Asn Tyr Phe Ile Asp Arg Asp Gly Pro Leu 50 55 60

Phe Arg Tyr Val Leu Asn Phe Leu Arg Thr Ser Glu Leu Thr Leu Pro 65 70 75 80

Leu Asp Phe Lys Glu Phe Asp Leu Leu Arg Lys Glu Ala Asp Phe Tyr 85 90 95

Gln Ile Glu Pro Leu Ile Gln Cys Leu Asn Asp Pro Lys Pro Leu Tyr 100 105 110

Pro Met Asp Thr Phe Glu Glu Val Val Glu Leu Ser Ser Thr Arg Lys 115 120 125

Leu Ser Lys Tyr Ser Asn Pro Val Ala Val Ile Ile Thr Gln Leu Thr 130 135 140

Ile Thr Thr Lys Val His Ser Leu Leu Glu Gly Ile Ser Asn Tyr Phe 145 150 155 160

Thr Lys Trp Asn Lys His Met Met Asp Thr Arg Asp Cys Gln Val Ser 165 170 175

Phe Thr Phe Gly Pro Cys Asp Tyr His Gln Glu Val Ser Leu Arg Val 180 185 190

His Leu Met Glu Tyr Ile Thr Lys Gln Gly Phe Thr Ile Arg Asn Thr 195 200 205

Arg Val His His Met Ser Glu Arg Ala Asn Glu Asn Thr Val Glu His 210 215 220

Asn Trp Thr Phe Cys Arg Leu Ala Arg Lys Thr Asp Asp 225 230 235

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Thr Leu Ala Thr Leu Leu Gln Asp Lys Ser Thr Leu Leu Ala Glu Leu 20 25 30

Phe Gly Glu Gly Arg Asp Ser Leu Ala Lys Asp Ser Lys Gly Arg Tyr 35 40 . 45

Phe Leu Asp Arg Asp Gly Val Leu Phe Arg Tyr Ile Leu Asp Phe Leu 50 55 60

Arg Asp Lys Ala Leu His Leu Pro Glu Gly Phe Arg Glu Arg Gln Arg 65 70 75 80

Leu Leu Arg Glu Ala Glu His Phe Lys Leu Thr Ala Met Leu Glu Cys 85 90 95

Ile Arg Ser Glu Arg Asp Ala Arg Pro Pro Gly Cys Ile Thr Ile Gly
100 105 110

Tyr Arg Gly Ser Phe Gln Phe Gly Lys Asp Gly Leu Ala Asp Val Lys 115 120 125

Phe Arg Lys Leu Ser Arg Ile Leu Val Cys Gly Arg Val Ala Gln Cys 130 135 140

Arg Glu Val Phe Gly Asp Thr Leu Asn Glu Ser Arg Asp Pro Asp His 145 150 155 160

Gly Gly Thr Asp Arg Tyr Thr Ser Arg Phe Phe Leu Lys His Cys Tyr 165 170 175

Ile Glu Gln Ala Phe Asp Asn Leu His Asp His Gly Tyr Arg Met Ala 180 185 190

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Gly Ser Cys Gly Ser Gly Thr Ala Gly Ser Ala Ala Glu Pro Lys Pro 195 200 205
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Gly Val Asp Thr Glu Glu Asn Arg Trp Asn His Tyr Asn Glu Phe Val 210 215 220

Phe Ile Arg Asp 225

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Tyr Thr Thr Thr Arg Ser Thr Leu Ser Lys Glu Thr Asp Thr Leu Leu 20 25 30

Ala Asn Ile Ala Ser Gly Ser Leu Ser Glu Asp Glu Gln Ala Asn Val 35 40 45

Val Thr Leu Pro Asp Gly Thr Leu Phe Val Asp Arg Asp Gly Pro Leu 50 55 60

Phe Ala Tyr Val Leu His Phe Leu Arg Thr Asp Lys Leu Ser Leu Pro 75 80

Glu Gln Phe Arg Glu Val Ala Arg Leu Lys Asp Glu Ala Asp Phe Tyr 85 90 95

Arg Leu Glu Arg Phe Ser Thr Leu Leu Ser Asn Ala Ser Ser Ile Ser 100 105 110

Pro Arg Pro Arg Thr Ala Asn Gly Tyr Asn Thr Ile Thr Ser Gly Ala 115 120 125

Glu Thr Gly Gly Tyr Ile Thr Leu Gly Tyr Arg Gly Thr Phe Ala Phe 130 135 140

Gly Arg Asp Gly Gln Ala Asp Val Lys Phe Arg Lys Leu His Arg Ile 145 150 155 160

Leu Val Cys Gly Arg Ala Thr Leu Cys Arg Glu Val Phe Ala Asp Thr 165 170 175

Leu Asn Glu Ser Arg Asp Pro Gly Gly Pro Asp Asp Gly Glu
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His Ile Asp Val Gly Gly His Met Tyr Thr Ser Ser Leu Ala Thr Leu 35 40 45

Thr Lys Tyr Pro Glu Ser Arg Ile Gly Arg Leu Phe Asp Gly Thr Glu 50 60

Pro Ile Val Leu Asp Ser Leu Lys Gln His Tyr Phe Ile Asp Arg Asp 65 70 75 80

Gly Gln Met Phe Arg Tyr Ile Leu Asn Phe Leu Arg Thr Ser Lys Leu 85 90 95

Leu Ile Pro Asp Asp Phe Lys Asp Tyr Thr Leu Leu Tyr Glu Glu Ala 100 105 110

Lys Tyr Phe Gln Leu Gln Pro Met Leu Leu Glu Met Glu Arg Trp Lys 115 120 125

Gln Asp Arg Glu Thr Gly Arg Phe Ser Arg Pro Cys Glu Cys Leu Val 130 135 140

Val Arg Val Ala Pro Asp Leu Gly Glu Arg Ile Thr Leu Ser Gly Asp 145 150 155 160

Lys Ser Leu Ile Glu Glu Val Phe Pro Glu Ile Gly Asp Val Met Cys 165 170 175

Asn Ser Val Asn Ala Gly Trp Asn His Asp Ser Thr His Val Ile Arg 180 185 190

Phe Pro Leu Asn Gly Tyr Cys His Leu Asn Ser Val Gln Val Leu Glu
195 200 205

Arg Leu Gln Gln Arg Gly Phe Glu Ile Val Gly Ser Cys Gly Gly 210 215 220

Val Asp Ser Ser Gln Phe Ser Glu Tyr Val Leu Arg Arg Glu Leu Arg 225 230 235 240

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Phe Phe Ile Asp Arg Asp Gly Phe Leu Phe Arg Tyr Ile Leu Asp Tyr
Leu Arg Asp Arg Gln Val Val Leu Pro Asp His Phe Pro Glu Lys Gly
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Arg Leu Lys Arg Glu Ala Glu Tyr Phe Gln Leu Pro Asp Leu Val Lys
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